# 510 (k) Summary - k081868

MAY 25 2011

Submitter

Department of the Army

U.S. Army Medical Research and Materiel Command

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**Date Summary Prepared** 

April 28, 2011

Trade/Proprietary Name

SMART Leish

Common or Usual Name

Leishmania Real-time Polymerase Chain Reaction

Diagnostic Assay,

Classification Name

Trypanosoma spp. serological reagents

**Regulation Number** 

866.3870

**Product Code** 

OUZ

**Review Panel** 

Microbiology 83

Classification

Class I

Predicate Device

Amizyme-Leishmania spp. Test Kit

### Intended Use

The **SMART Leish** is a qualitative diagnostic real-time PCR test for the detection of *Leishmania* species and the identification of *L. major* in skin lesion scrapings and punch biopsies from individuals suspected of having cutaneous leishmaniasis. The test utilizes real-time polymerase chain reaction assay on the Cepheid SmartCycler® II Dx to detect *Leishmania* species and *L. major*.

### WARNING

The **SMART Leish** is intended for use only by trained laboratory personnel in Department of Defense laboratories. Clinical performance has not been established with strains other than *L. major*.

# **Device Description**

The **SMART Leish** consists of an assay reagent kit and assay definition files for the polymerase chain reaction (PCR) instrument platform. The kit contains sufficient reagents, in lyophilized bead form, to qualitatively assay 50 clinical samples for both *Leishmania* genus and *L. major* targets. Additional required accessories that are specified include the PCR instrument platform, a deoxyribonucleic acid (DNA) purification kit, and a positive extraction control. The device components and required accessories are listed as follows.

For the Leishmania assays, a tissue specimen (skin scraping or punch biopsy) from an individual suspected of being infected with Leishmania species or L. major is collected in 70% or 100% ethanol, and the DNA is extracted from the specimen using the Qiagen QIAamp DNA Mini Kit. An aliquot of this DNA is tested using the Leishmania genus assay, which will amplify a portion of DNA encoding the Leishmania species 16S ribosomal ribonucleic acid (rRNA) gene if present. Amplified targets are detected using a TagMan® hybridization probe with 6-carboxyfluorescein.(FAM) reporter dye (517 nm) and a Black Hole Quencher® (BHQ). This assay also contains a positive internal control consisting of a nonsensical, non-naturally occurring DNA sequence, with Texas Red reporter dye (615 nm) and BHQ, used to detect evidence of PCR inhibition and confirm the integrity of assay reagents in negative specimens. If the sample tests positive for Leishmania species, another aliquot of DNA may be tested using the L. major assay, which will amplify a portion of DNA encoding the GPI gene that is specific to L. major. Amplified targets are detected using a TaqMan hybridization probe with FAM reporter dye and a BHQ. The DNA amplification, detection of fluorescence, and interpretation of signals are done automatically by the SmartCycler instrument for each assay. The thermocycling protocols take approximately 45-55 minutes.

# Substantial Equivalence Information

- Predicate device name:
   Amizyme-Leishmania spp. Test Kit
- 2. 510k number : K842526

# Comparison with predicate:

Table 1. Similarities

Device Aspect	SMART Leish	Reference Device		
Intended Use	Detection of Leishmania spp	Detection of Leishmania spp		
Assay Type	Qualitative	Qualitative		
Agent measured	Leishmania genus and L. major DNA	Leishmania spp		
Technology	Qiagen QIAap DNA Mini Kit	None		
Controls	External – positive and negative controls for both the extraction process and detection assay runs  Internal – positive control for PCR inhibitors in sample	External positive and negative controls		
	Differences			
Instrument Platform	Cepheid SmartCycler	None		
Automation	Automated DNA amplification cycling, probe detection, and interpretation of results	Manual method		
Sample Collection	Skin punch biopsy or scrapings, stored in ethanol	Serum		
Endpoint detection method	PCR amplification of DNA sequences unique to the organisms of interest with	Indirect Immunofluoresence (IFA) Fluorescent conjugate secondary		

# Clinical Performance Testing

The clinical performance of the **SMART** Leish assay was established in a multi-center, retrospective clinical study conducted at two military U.S. hospital sites.

A total of 312 Leishmania genus and 187 Leishmania major prospective specimens were collected from adults 18 years or older and evaluated in the SMART Leish assay and compared to culture and/or microscopy. Evaluated specimens were skin lesion scrapings and punch biopsies from individuals suspected of having cutaneous leishmaniasis. The patient population consisted entirely of military personnel who were at the time, or had previously been, deployed to Southwest Asia—primarily Afghanistan and Iraq. SMART Leish performance versus culture and microcopy, including 95% confidence intervals, is detailed below.

Table 2. Summary of SMART Leish Combined Studies for the Leishmania genus Assay

		Clinical Truth			
		Positive	Negative	Total	
SMART Leish -	Positive	223	29	252	
Leishmania Genus Assay	Negative	2	58	60	
	Total	225	87	312	

<sup>\*</sup>Diagnosis of leishmaniasis by culture and slide techniques results in lower clinical sensitivity than PCR assays. In this clinical data set there were 29 false positives generated for the *Leishmania* Genus assay. Twenty seven of these Genus false positives were determined to be Genus positive using sequencing methods.

Table 3. Summary of SMART Leish Combined Studies for the Leishmania major Assay

		Clinical Truth			
		Positive	Negative	Total	
SMART Leish - L.	Positive	92	8	100	
major Assay	Negative	4	83	87	
	Total	96	91	187	

Diagnosis of leishmaniasis by culture and slide techniques results in lower clinical sensitivity than PCR assays. In this clinical data set there were eight false positives generated for the *L. major* assay, of which three of were determined to be *L. major* using sequencing methods.

# **ANALYTICAL PERFORMANCE**

### **Analytical Reactivity Study**

Two separate studies were done with Leishmania genus and Leishmania major assays using 36 Leishmania strains. The goal was to determine whether the assays cross-reacts with the nucleic acids from non-target organisms. In the Leishmania genus study, DNA from 36 *Leishmania* strains was tested and gave robust positive results with the Leishmania genus PCR assay. In the Leishmania major study, 11 of 36 strains were Leishmania major species. All 11 tested gave robust positive results with the *Leishmania major* PCR assay. None of the 25 non-*Leishmania major* samples tested positive. See **Table 4**.

Table 4. List of Leishmania Organisms Tested in Reactivity Study

Leishmania major (11)	Leishmania mexicana
Leishmania tropica (9)	Leishmania amazonensis
Leishmania infantum (7)	Leishmania peruviana
Leishmania donovani (2)	Leishmania guyanensis (2)
Leishmania braziliensis	Leishmania tarentolae

#### **Analytical Specificity**

A total of 121 DNA samples were evaluated in the specificity study. The DNA from 11 *Leishmania major* strains tested in this study gave robust positive results with the *Leishmania major* PCR assay. Twenty-five non-*Leishmania major* samples tested negative; hence the analytical specificity was 100%. The percentage of false positives was 0%, and the percentage of false negatives is 0%. In the *Leishmania genus* study, 85 of the 121 samples were non-target DNA samples. Eighty-five were negative, 2 false positives and 36 were positive. The analytical specificity of the *Leishmania genus* PCR assay was 98.3%, false positive rate was 2.4% and the false negative rate was 0%. See **Table 5** for list of organisms used in the study.

Nucleic acids from non-target organisms were tested for both the *Leishmania* genus and *Leishmania major* assays. DNA samples with concentrations used in this study included purified DNA from bacteria (52; conc =50 - 46,250 pg/uL), fungi (11; conc= 50,000 - 97,000 pg/uL), viruses (7; conc= 5,950 - 30,990 pg/uL), mammals (3; conc = 20 pg/uL (human, bovine, and murine)), human melanoma cell lines (3; conc = 10,200 - 25,200 pg/uL), *Leishmania major* (11; conc = 21,540 - 83,000 pg/uL), 25 additional *Leishmania* species (not-*L. major*) (conc = 16,680 - 592,00 pg/uL), and 9 trypanosomes representing 6 species (conc = 12,760 - 95,000 pg/uL).

Table 5. List of Organisms Used in Specificity Study

	Bacteria
	Acinetobacter baumanni
	Bacillus anthracis (3)
	Bacillus cereus (2)
	Bacillus subtilis var. niger
	Bacillus thuringiensis
	Brucella abortus
	Brucella melitiensis
	Brucella suis
	Clostridium botulinum type A
	Clostridium botulinum type B
	Clostridium perfringens (2)
İ	Clostridium sordelli
	Enterobacter aerogenes
İ	Enterococcus durans
1	Enterococcus faecalis
	Escherichia coli
	Francisella tularensis (2)
	Haemophilus influenzae
	Klebsiella oxytoca
	Klebsiella pneumoniae
	Moraxella cattaharalis
	Neisseria lactamica
	Pasteurella multocida
	Proteus mirabilis
İ	Proteus vulgaris
	Providencia stuartii

Pseudomonas aeruginosa Staphylococcus aureus (4) Staphylococcus hominis Stenotrophomonas maltophilia Streptococcus pyogenes (2) Streptococcus sp. (B) Streptococcus (F2) Yesinia pestis Yersinia

# Mycobacteria Mycobacteria abscessus Mycobacteria fortuitum Mycobacteria marinum Mycobacteria spp Mycobacteria tuberculosis

Mycobacteria ulcerans

Viruses Human Papilloma Virus Herpes Simplex Virus Type I Herpes Simplex Virus Type II (2) Varicella Zoster Virus (2)

Fungi Microsporium gypseum Cladophialophora carrionii Fonsecaea pedrosoi Rhinocladiella compacta
Phialophora verrucosa
Trichophyton tonsurans
Trichophyton mentagrophytes
Trichophyton soudanense
Arthroderma benhamiae
Sporothrix schenckii
Microsporum canis

### Mammalian Bovine Human Murine Human melanoma cell line (3)

Trypanosoma
Trypanosoma cruzi (2)
Trypanosoma rhodesiense
Trypanosoma rangeli (2)
Trypanosomal lewisi lincicome
Trypanosoma brucei gambiense
Crithidia fasciculata (2)

Two strains of *C. fasciculata* were determined to be low-level cross-reactors based on false-positive results observed. The degree of cross-reactivity of the *Leishmania* genus assay for *C. fasciculata* can be considered low level or weak. No cross-reactors were determined for the *L. major* assay through the study.

Supplementary *in silico* analysis was done comparing the Smart Leish primer and probe sequences to sequences from the following organisms that could potentially result in a clinical presentation similar to cutaneous leishmaniasis: Corynebacterium diphtheria (veldt sores), Mycobacteria (Tropical Ulcer), Mycobacterium tuberculosis (Lupus vulgaris), Treponema pallidum (Tertiary Syphilis), Treponema pertenue (Yaws), and Blastomyces (Blastomycosis). Each Smart Leish primer and probe contained at least four mismatches to those sequences. Based on these alignments, no cross-reactivity was predicted with the Smart Leish primers and probes.

# **Analytical Sensitivity**

The assay linearity of SMART Leish for *Leishmainia* genus and *Leishmania major* was determined using eight different concentrations of purified *L. major* DNA tested in a 10-fold dilution series from 14.26 ng to 0.001426 pg and 21.39 ng to 0.002139 pg, respectively, DNA per reaction. Based on these results, additional studies were done to determine the analytical sensitivity (limit of detection or LOD) which was defined as the lowest amount of purified *L. major* DNA that consistently provided positive test results 95% of the time.

In addition to the LOD determination, six different concentrations of purified L. major parasites were tested in a 10-fold dilution of 5 X  $10^2$  parasites/mL (equivalent to  $10^7$  to  $10^2$  parasites/extraction) to determine the minimum number of L. major parasites that can be taken through the extraction and testing procedure and still give a positive result for the SMART Leish assays (extraction LOD).

Table 6 summarizes the results of the assay LOD for the *L. genus* and *L. major* assays, and Table 7 summarizes the results of the extraction LOD (eLOD) study.

Table 6. Summary of the Measuring Range and LOD for the *Leishmania* genus and *L. major* Assays (Values are per assay reaction)

Assay	Linear Range <sup>1</sup>	Assay LOD <sup>1,2</sup>	LOD Equates to
Leishmania Genus Assay	0.014–14,260 pg	0.14 pg	4 genome copies with ~200 copies/genome o target gene
L. major Assay	0.2139–21,390 pg	2.1 pg	62 genome copies

<sup>&</sup>lt;sup>1</sup>Amount of *L. major* DNA in the PCR reaction. LOD refers to the minimum amount of L. major DNA that can be put into a SMART Leish PCR reaction and still result in a positive SMART Leish test greater than or equal to 95% of the time.

Table 7. Summary of Assay eLOD for the *Leishmania* genus and *L. major* Assays Using Parasite Samples. Values are per extraction.

Assay	eLOD <sup>1</sup>	LOD Equates to
Leishmania Genus Assay	250 parasites	~5 genome copies at assay reaction stage
L. major Assay	1,000 parasites	~30 genome copies at assay reaction stage

<sup>195%</sup> accuracy across different operators and days and from multiple L. major strains

### Reproducibility

Precision was evaluated internally at the study sites. The following tables summarize the between laboratory reproducibility results for the *Leishmania* genus and *L. major* assays, respectively. These tables include the results of the studies done with purified DNA, cultured parasites, and mock human samples. The negative mock human samples were not further tested using the *L. major* assay if the initial *Leishmania* genus assay result was negative.

<sup>&</sup>lt;sup>2</sup>95% accuracy across different operators and instruments and from multiple L. major strains

Extraction LOD refers to the minimum number of L. major parasites that can be taken through the Qiagen extraction procedure
and still consistently result in a positive SMART Leish test greater than or equal to 95% of the time. Extraction LOD was
determined across different operators, different days, and using multiple L. major strains.

Table 8. Summary of Between Laboratory Reproducibility Results for the *Leishmania* Genus Assay

Sample Type	Sample ID	Concentration	Number of Replicates (over 5 days)	Laboratory A (WRAIR)	Laboratory B (BAMC)	Laboratory C (MAMC)	Total Agreemen t	% Total Agreemen t
		genome equivalents/µL						
Purified DNA	Low	2.0	20	20/20	20/20	20/20	60/60	100%
	Medium	20	20	20/20	20/20	20/20	60/60	100%
	High	200	20	20/20	20/20	20/20	60/60	100%
Positive Agreement for Purified DNA							180/180	100%
		(parasites/						
Cultured Parasites	Low	1,000	20	20/20	20/20	20/20	60/60	100%
	Medium	10,000	20	20/20	20/20	20/20	60/60	100%
	High	25,000	20	20/20	20/20	20/20	60/60	100%
Positive Agreement for Cultured Parasites							180/180	100%
		(parasites/	,					
Mock Human Samples	Negative	0	10	10/10	10/10	10/10	30/30	100%
	Positive	1.1x10^6	14	14/14	13/14	14/14	41/42	97.6%
Overall Agreement for Mock Human Samples	2						71/72	98.6%

Table 9. Summary of between Laboratory Reproducibility Results for the Leishmania major Assay

Sample Type	Sample ID	Concentration	Number of Replicates (over 5 days)	Laboratory A (WRAIR)	Laboratory B (BAMC)	Laboratory C (MAMC)	Total Agreem ent	% Total Agreement
		genome equivalents/µL						
Purified DNA	Low	2.0	20	15/20	7/20	15/20	37/60	61%
	Medium	20	20	20/20	20/20	20/20	60/60	100%
	High	200	20	20/20	20/20	20/20	60/60	100%
Positive Agreement for Purified DNA							157/180	87.2%
		(parasites/ extraction)						
Cultured Parasites	Low	1,000	20	19/20	20/20	16/20	55/60	91.6%
	Medium	10,000	20	20/20	20/20	20/20	60/60	100%
	High	25,000	20	20/20	20/20	20/20	60/60	100%
Positive Agreement for Cultured Parasites						·	175/180	97.2% CI
		(parasites/						
Mock Human Samples	Negative	0	0/0	0/0	0/0	0/0	0/0	100%
	Positive	1.1x10^6	14	14/14	13/13	14/14	41/41	100%
Overall Agreement for Mock Human Samples							41/41	100%

### INTERFERING SUBSTANCES

The *Leishmania* genus assay and the multiplexed positive IC assay were evaluated with known PCR inhibitors. The inhibitors tested were: hemoglobin, hemin, ferric ammonium citrate (FAC), and EDTA. Presence of inhibition was defined as a false negative result (negative result for a true positive sample). The inhibitor concentrations tested and the maximum inhibitor concentration that did not induce inhibition in the *Leishmania* genus and positive IC assays are listed in the table below.

Table 10. Results of Testing the Effects of Known Inhibitors on the Leishmania Genus and IC Assays

PCR Inhibitor	Concentrations Tested (final concentration in PCR	Maximum Concentration without Inhibition Detected		
	reaction tube)	Leishmania Genus Assay	IC Assay	
Hemoglobin	0.03, 0.06, 0.12, 0.24, 0.48, 0.96, and 1.92 µg/µl	1.92 µg/µl	0.96 µg/µl	
Hemin	0.03, 0.06, 0.12, 0.24, 0.48, 0.96, and 1.92 µg/µl	0.96 µg/µl	0.96 µg/µl	
FAC	0.03125, 0.0625, 0.125, 0.25, 0.50, and 1.00%	0.0625%	0.0625%	
EDTA	0.3125, 0.625, 1.25, 2.5, 5, and 10 mM	1.25 mM	1.25 mM	

# Conclusion from Nonclinical and Clinical Performance Testing

The conclusions drawn from the nonclinical and clinical testing, and comparative methodology review, demonstrate the effectiveness of the **SMART Leish** for the intended use.

## DEPARTMENT OF HEALTH & HUMAN SERVICES



MAY 2 5 2011



Food and Drug Administration 10903 New Hampshire Avenue Silver Spring, MD 20993

Department of the Army
U.S Army Medical Material Development Activity
c/o Robert E. Miller, Ph.D., RAC
Director, Division of Regulated Activities and Compliance
1430 Veterans Drive
Fort Detrick, MD 21702-5009

Re: k081868

Trade/Device Name: SMART Leish PCR Assay

Regulation Number: 21 CFR 866.3870

Regulation Name: Trypanosoma spp. Serological Reagents

Regulatory Class: Class I

Product Code: OUZ Dated: May 23, 2011 Received: May 24, 2011

Dear Dr. Miller:

We have reviewed your Section 510(k) premarket notification of intent to market the device referenced above and have determined the device is substantially equivalent (for the indications for use stated in the enclosure) to legally marketed predicate devices marketed in interstate commerce prior to May 28, 1976, the enactment date of the Medical Device Amendments, or to devices that have been reclassified in accordance with the provisions of the Federal Food, Drug, and Cosmetic Act (Act) that do not require approval of a premarket approval application (PMA). You may, therefore, market the device, subject to the general controls provisions of the Act. The general controls provisions of the Act include requirements for annual registration, listing of devices, good manufacturing practice, labeling, and prohibitions against misbranding and adulteration.

If your device is classified (see above) into class II (Special Controls), it may be subject to such additional controls. Existing major regulations affecting your device can be found in Title 21, Code of Federal Regulations (CFR), Parts 800 to 895. In addition, FDA may publish further announcements concerning your device in the Federal Register.

Please be advised that FDA's issuance of a substantial equivalence determination does not mean that FDA has made a determination that your device complies with other requirements of the Act or any Federal statutes and regulations administered by other Federal agencies. You must comply with all the Act's requirements, including, but not limited to: registration and listing (21 CFR Part 807); labeling (21 CFR Parts 801 and 809); medical device reporting (reporting of medical device-related adverse events) (21 CFR 803); and good manufacturing practice requirements as set forth in the quality systems (QS) regulation (21 CFR Part 820). This letter will allow you to begin marketing your device as described in your Section 510(k) premarket

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notification. The FDA finding of substantial equivalence of your device to a legally marketed predicate device results in a classification for your device and thus, permits your device to proceed to the market.

If you desire specific advice for your device on our labeling regulation (21 CFR Parts 801 and 809), please contact the Office of In Vitro Diagnostic Device Evaluation and Safety at (301) 796-5450. Also, please note the regulation entitled, "Misbranding by reference to premarket notification" (21 CFR Part 807.97). For questions regarding the reporting of adverse events under the MDR regulation (21 CFR Part 803), please go to http://www.fda.gov/MedicalDevices/Safety/ReportaProblem/default.htm for the CDRH's Office of Surveillance and Biometrics/Division of Postmarket Surveillance.

You may obtain other general information on your responsibilities under the Act from the Division of Small Manufacturers, International and Consumer Assistance at its toll-free number (800) 638-2041 or (301) 796-7100 or at its Internet address http://www.fda.gov/cdrh/industry/support/index.html.

Sincerely yours,

Sally A. Hojvat, Ph.D., MSc.

Director

Division of Microbiology Devices

Office of *In Vitro* Diagnostic Device Evaluation and Safety

Center for Devices and Radiological Health

# INDICATIONS FOR USE STATEMENT

510(k) Number (if known): <u>k081868</u>

Device Name:

SMART Leish

Indications for Use:

The **SMART Leish** is a qualitative diagnostic real-time PCR test for the rapid detection of *Leishmania* species and the identification of *L. major* in skin lesion scrapings and punch biopsies from individuals suspected of having cutaneous leishmaniasis. The test utilizes real-time polymerase chain reaction assay on the Cepheid SmartCycler II Dx to detect *Leishmania* species and *L. major*. The **SMART Leish** is indicated for use in patients with clinical presentations and travel history.

### WARNING

The **SMART Leish** is intended for use only by trained laboratory personnel in Department of Defense laboratories. Clinical performance has not been established with strains other than *L. major*.

Prescription Use 
√ AND/OR Over-The-Counter Use \_\_\_\_\_
(Part 21 CFR 801 Subpart D) (21 CFR 801 Subpart C)
(PLEASE DO NOT WRITE BELOW THIS LINE-CONTINUE ON ANOTHER PAGE OF NEEDED)

Concurrence of CDRH, Office of In Vitro Diagnostic Devices (OIVD)

Division Sign-Off

Office of In Vitro Diagnostic Device Evaluation and Safety

SMART Leish 510(k) K081867

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510(k) Notification